RAW SEQUENCE LISTING   DATE: 01/15/2002
PATENT APPLICATION: US/09/990,046   TIME: 18:55:44

Input Set : **N:\Crf3\RULE60\09990046.raw**
Output Set: **N:\CRF3\01152002\I990046.raw**

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 1 <110> APPLICANT: de Sauvage, Frederic
 2        Carpenter, David A.
 3 <120> TITLE OF INVENTION: Patched-2
 4 <130> FILE REFERENCE: P1405R1
 5 <140> CURRENT APPLICATION NUMBER: US/09/990,046
 6 <141> CURRENT FILING DATE: 2001-11-20
 7 <150> PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/293,505
 8 <151> PRIOR FILING DATE: EARLIER FILING DATE: 1999-04-15
 9 <160> NUMBER OF SEQ ID NOS: 32
11 <210> SEQ ID NO: 1
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13 <212> TYPE: DNA
14 <213> ORGANISM: Homo sapiens
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18        aaaatgcttt atttgtgaaa tttgtgatgc tattgcttta tttgtaacca 150
19        ttataagctg caataaacaa gttgggccat ggcggccaag cttctgcagg 200
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21        ctcagagagc tgcccccgag ttacacaccc ccagctcgaa ccgcagcacc 300
22        ccagatccta gctgggagcc tgaaggctcc actctggctt cgtgcttact 350
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24        aaagtgctct ttctgggact gttggccttt ggggccctgg cattaggtct 450
25        ccgcatggcc attattgaga caaacttgga acagctctgg gtagaagtgg 500
26        gcagccgggt gagccaggag ctgcattaca ccaaggagaa gctgggggag 550
27        gaggctgcat acacctctca gatgctgata cagaccgcac gccaggaggg 600
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30        aacaaaatct gctacaagtc aggagttccc cttattgaaa atggaatgat 750
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57        gaagccagca gccagcatgt ggtcaccatc ctgcctcccc aagcccacct 2100
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59        ggtccacacg ggaccttcta ggccaggagg aggagacaag gcagaaggca 2200
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98 <210> SEQ ID NO: 2
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PATENT APPLICATION: US/09/990,046   TIME: 18:55:44

Input Set : N:\Crf3\RULE60\09990046.raw
Output Set: N:\CRF3\01152002\I990046.raw

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101 <213> ORGANISM: Homo sapiens
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107         Lys Ala Pro Leu Trp Leu Arg Ala Tyr Phe Gln Gly Leu Leu Phe
108                         35                  40                  45
109         Ser Leu Gly Cys Gly Ile Gln Arg His Cys Gly Lys Val Leu Phe
110                         50                  55                  60
111         Leu Gly Leu Leu Ala Phe Gly Ala Leu Ala Leu Gly Leu Arg Met
112                         65                  70                  75
113         Ala Ile Ile Glu Thr Asn Leu Glu Gln Leu Trp Val Glu Val Gly
114                         80                  85                  90
115         Ser Arg Val Ser Gln Glu Leu His Tyr Thr Lys Glu Lys Leu Gly
116                         95                  100                 105
117         Glu Glu Ala Ala Tyr Thr Ser Gln Met Leu Ile Gln Thr Ala Arg
118                         110                 115                 120
119         Gln Glu Gly Glu Asn Ile Leu Thr Pro Glu Ala Leu Gly Leu His
120                         125                 130                 135
121         Leu Gln Ala Ala Leu Thr Ala Ser Lys Val Gln Val Ser Leu Tyr
122                         140                 145                 150
123         Gly Lys Ser Trp Asp Leu Asn Lys Ile Cys Tyr Lys Ser Gly Val
124                         155                 160                 165
125         Pro Leu Ile Glu Asn Gly Met Ile Glu Trp Met Ile Glu Lys Leu
126                         170                 175                 180
127         Phe Pro Cys Val Ile Leu Thr Pro Leu Asp Cys Phe Trp Glu Gly
128                         185                 190                 195
129         Ala Lys Leu Gln Gly Gly Ser Ala Tyr Leu Pro Gly Arg Pro Asp
130                         200                 205                 210
131         Ile Gln Trp Thr Asn Leu Asp Pro Glu Gln Leu Leu Glu Glu Leu
132                         215                 220                 225
133         Gly Pro Phe Ala Ser Leu Glu Gly Phe Arg Glu Leu Leu Asp Lys
134                         230                 235                 240
135         Ala Gln Val Gly Gln Ala Tyr Val Gly Arg Pro Cys Leu His Pro
136                         245                 250                 255
137         Asp Asp Leu His Cys Pro Pro Ser Ala Pro Asn His His Ser Arg
138                         260                 265                 270
139         Gln Ala Pro Asn Val Ala His Glu Leu Ser Gly Gly Cys His Gly
140                         275                 280                 285
141         Phe Ser His Lys Phe Met His Trp Gln Glu Glu Leu Leu Leu Gly
142                         290                 295                 300
143         Gly Met Ala Arg Asp Pro Gln Gly Glu Leu Leu Arg Ala Glu Ala
144                         305                 310                 315
145         Leu Gln Ser Thr Phe Leu Leu Met Ser Pro Arg Gln Leu Tyr Glu
146                         320                 325                 330
147         His Phe Arg Gly Asp Tyr Gln Thr His Asp Ile Gly Trp Ser Glu
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Output Set: N:\CRF3\01152002\I990046.raw

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149     Glu Gln Ala Ser Thr Val Leu Gln Ala Trp Gln Arg Arg Phe Val
150                     350                 355                 360
151     Gln Leu Ala Gln Glu Ala Leu Pro Glu Asn Ala Ser Gln Gln Ile
152                     365                 370                 375
153     His Ala Phe Ser Ser Thr Thr Leu Asp Asp Ile Leu His Ala Phe
154                     380                 385                 390
155     Ser Glu Val Ser Ala Ala Arg Val Val Gly Gly Tyr Leu Leu Met
156                     395                 400                 405
157     Leu Ala Tyr Ala Cys Val Thr Met Leu Arg Trp Asp Cys Ala Gln
158                     410                 415                 420
159     Ser Gln Gly Ser Val Gly Leu Ala Gly Val Leu Leu Val Ala Leu
160                     425                 430                 435
161     Ala Val Ala Ser Gly Leu Gly Leu Cys Ala Leu Leu Gly Ile Thr
162                     440                 445                 450
163     Phe Asn Ala Ala Thr Thr Gln Val Leu Pro Phe Leu Ala Leu Gly
164                     455                 460                 465
165     Ile Gly Val Asp Asp Val Phe Leu Leu Ala His Ala Phe Thr Glu
166                     470                 475                 480
167     Ala Leu Pro Gly Thr Pro Leu Gln Glu Arg Met Gly Glu Cys Leu
168                     485                 490                 495
169     Gln Arg Thr Gly Thr Ser Val Val Leu Thr Ser Ile Asn Asn Met
170                     500                 505                 510
171     Ala Ala Phe Leu Met Ala Ala Leu Val Pro Ile Pro Ala Leu Arg
172                     515                 520                 525
173     Ala Phe Ser Leu Gln Ala Ala Ile Val Val Gly Cys Thr Phe Val
174                     530                 535                 540
175     Ala Val Met Leu Val Phe Pro Ala Ile Leu Ser Leu Asp Leu Arg
176                     545                 550                 555
177     Arg Arg His Cys Gln Arg Leu Asp Val Leu Cys Cys Phe Ser Ser
178                     560                 565                 570
179     Pro Cys Ser Ala Gln Val Ile Gln Ile Leu Pro Gln Glu Leu Gly
180                     575                 580                 585
181     Asp Gly Thr Val Pro Val Gly Ile Ala His Leu Thr Ala Thr Val
182                     590                 595                 600
183     Gln Ala Phe Thr His Cys Glu Ala Ser Ser Gln His Val Val Thr
184                     605                 610                 615
185     Ile Leu Pro Pro Gln Ala His Leu Val Pro Pro Pro Ser Asp Pro
186                     620                 625                 630
187     Leu Gly Ser Glu Leu Phe Ser Pro Gly Gly Ser Thr Arg Asp Leu
188                     635                 640                 645
189     Leu Gly Gln Glu Glu Glu Thr Arg Gln Lys Ala Ala Cys Lys Ser
190                     650                 655                 660
191     Leu Pro Cys Ala Arg Trp Asn Leu Ala His Phe Ala Arg Tyr Gln
192                     665                 670                 675
193     Phe Ala Pro Leu Leu Leu Gln Ser His Ala Lys Ala Ile Val Leu
194                     680                 685                 690
195     Val Leu Phe Gly Ala Leu Leu Gly Leu Ser Leu Tyr Gly Ala Thr
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Input Set : **N:\Crf3\RULE60\09990046.raw**
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197	Leu Val Gln Asp Gly Leu Ala Leu Thr Asp Val Val Pro Arg Gly	
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199	Thr Lys Glu His Ala Phe Leu Ser Ala Gln Leu Arg Tyr Phe Ser	
200	725 730 735	
201	Leu Tyr Glu Val Ala Leu Val Thr Gln Gly Gly Phe Asp Tyr Ala	
202	740 745 750	
203	His Ser Gln Arg Ala Leu Phe Asp Leu His Gln Arg Phe Ser Ser	
204	755 760 765	
205	Leu Lys Ala Val Leu Pro Pro Pro Ala Thr Gln Ala Pro Arg Thr	
206	770 775 780	
207	Trp Leu His Tyr Tyr Arg Asn Trp Leu Gln Gly Ile Gln Ala Ala	
208	785 790 795	
209	Phe Asp Gln Asp Trp Ala Ser Gly Arg Ile Thr Arg His Ser Tyr	
210	800 805 810	
211	Arg Asn Gly Ser Glu Asp Gly Ala Leu Ala Tyr Lys Leu Leu Ile	
212	815 820 825	
213	Gln Thr Gly Asp Ala Gln Glu Pro Leu Asp Phe Ser Gln Leu Thr	
214	830 835 840	
215	Thr Arg Lys Leu Val Asp Arg Glu Gly Leu Ile Pro Pro Glu Leu	
216	845 850 855	
217	Phe Tyr Met Gly Leu Thr Val Trp Val Ser Ser Asp Pro Leu Gly	
218	860 865 870	
219	Leu Ala Ala Ser Gln Ala Asn Phe Tyr Pro Pro Pro Glu Trp	
220	875 880 885	
221	Leu His Asp Lys Tyr Asp Thr Thr Gly Glu Asn Leu Arg Ile Pro	
222	890 895 900	
223	Pro Ala Gln Pro Leu Glu Phe Ala Gln Phe Pro Phe Leu Leu Arg	
224	905 910 915	
225	Gly Leu Gln Lys Thr Ala Asp Phe Val Glu Ala Ile Glu Gly Ala	
226	920 925 930	
227	Arg Ala Ala Cys Ala Glu Ala Gly Gln Ala Gly Val His Ala Tyr	
228	935 940 945	
229	Pro Ser Gly Ser Pro Phe Leu Phe Trp Glu Gln Tyr Leu Gly Leu	
230	950 955 960	
231	Arg Arg Cys Phe Leu Leu Ala Val Cys Ile Leu Leu Val Cys Thr	
232	965 970 975	
233	Phe Leu Val Cys Ala Leu Leu Leu Leu Asn Pro Trp Thr Ala Gly	
234	980 985 990	
235	Leu Ile Val Leu Val Leu Ala Met Met Thr Val Glu Leu Phe Gly	
236	995 1000 1005	
237	Ile Met Gly Phe Leu Gly Ile Lys Leu Ser Ala Ile Pro Val Val	
238	1010 1015 1020	
239	Ile Leu Val Ala Ser Val Gly Ile Gly Val Glu Phe Thr Val His	
240	1025 1030 1035	
241	Val Ala Leu Gly Phe Leu Thr Thr Gln Gly Ser Arg Asn Leu Arg	
242	1040 1045 1050	
243	Ala Ala His Ala Leu Glu His Thr Phe Ala Pro Val Thr Asp Gly	
244	1055 1060 1065	
245	Ala Ile Ser Thr Leu Leu Gly Leu Leu Met Leu Ala Gly Ser His	

VERIFICATION SUMMARY   DATE: 01/15/2002
PATENT APPLICATION: **US/09/990,046**   TIME: 18:55:45

Input Set : **N:\Crf3\RULE60\09990046.raw**
Output Set: **N:\CRF3\01152002\I990046.raw**

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L:275 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:277 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:278 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:279 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:304 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:5
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